PTORSAGBA(08-03)
Approved for use through 07/31/2009, OMB 0851-0031
Us Pater 8 Trademont Ottoc: U.S. DEPARTMENT OF CONDUCTED
ter the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid CMB control number.
Complete if Known Substitute for form 1449APTO INFORMATION DISCLOSURE **Application Number** 09/687,483 STATEMENT BY APPLICANT October 13, 2000 Filing Date BRAUN, Andreas, et al. First Named Inventor 1631 **Group Art Unit** CLOW, Lori A **Examiner Name** Attorney Docket No: SEQ-2033-CP

	A YHADE	US PA	ATENT DOCUMENT	
Examiner Initial *	USP Document Number	Publication Date	Name of Patentee or Applicant of cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear

		FOREIGN PATE	ENT DOCUMENTS		
Examiner Initials*	Foreign Document No	Publication Date	Name of Patentee or Applicant of cited Document	Pages,Columns,Lines, Where Relevant Passages or Relevant Figures Appear	T²

	OTHE	R DOCUMENTS NON PATENT LITERATURE DOCUMENTS	
Examiner Initials*	Cite No 1	include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	
LACT		BREEN, G., et al., Determining SNP Allele Frequencies in DNA Pools, BioTechniques, (2000), 464-470, 28(3).	
		BUETOW, Kenneth H. et al., High-throughput development and characterization of a genomewide collection of gene-based single nucleotide polymorphism markers by chip-based matrix-assisted laser desorption/ionization time-of-flight mass spectrometry, Proc. National Association of Science (2001), 581-584, 98(2) PNAS http://www.pnas.org .	
		DOWNES, Kate, et al., SNP allele frequency estimation in DNA pools and variance components analysis, BioTechniques, (2004), 840-845, 36(5), The Wellcome Trust Sanger Institute.	
	<u> </u>	GERMER, Soren, et al., High-Throughput SNP Allele-Frequency Determination in Pooled DNA Samples by Kinetic PCR, Methods, Genome Research, (2000), 258-266, 10, Cold Spring Harbor Laboratory Press.	
		HOOGENDOORN, Bastiaan, et al., Cheap, accurate and rapid allele frequency estimation of single nucleotide polymorphisms by primer extension and DHPLC in DNA pools, Hum Genet (2000) 488-493, 107, Pringer-Verlag.	
		LAKEN, Steven J. et al., Genotyping by mass spectrometric analysis of short DNA fragments, Research, Nature Biotechnology, (1998), 1352-1356, 16, Nature America Inc. (http://biotech.nature.com).	
,		LE HELLARD, Stephanie, et al., SNP genotyping on pooled DNA's: comparison of genotyping technologies and a semi automated method for data storage and analysis, Nucleic Acids Research, (2002) 1-10, 30(15), Oxford University Press.	
		RISH, Neil, et al., The Relative Power of Family-Based and Case-Control Designs for Linkage Disequilibrium Studies of Complex Human Diseases I. DNA Pooling, Genome Research, (1998), 1273-1288, 8, Cold Spring Harbor Laboratory Press.	-
4		ROSS, Philip, et al., Quantitative Approach to Single-Nucleotide Polymorphism Analysis Using MALDI-TOF Mass Spectrometry, BioTechniques, (2000) 620-629, 29(3).	
LAC		SASAKI, Tomonari, et al., Precise Estimation of Allele Frequencies of Single-Nucleotide Polymorphisms by a Quantitative SSCP Analysis of Pooled DNA, Am. J. Hum, Genet. (2001)	

EXAMINER	Lou' A-Claw	DATE CONSIDERED	3/16	105
		والمستحدد		

PTO/SE/08A(08-03

Approved for use through 07/31/2008, CM/8 0651-003

Substitute for form 1449A/PTO INFORMATION DISCLOSURE **Application Number** 09/687,483 STATEMENT BY APPLICANTE October 13, 2000 **Filing Date First Named Inventor** BRAUN, Andreas, et al. MAR 0 2 2005 1631 **Group Art Unit** CLOW, Lori A **Examiner Name** Attorney Docket No: SEQ-2033-CP Sheet 2 of 2

	OTHE	R DOCUMENTS - NON PATENT LITERATURE DOCUMENTS	
Examiner Initials*	Cite No 1	include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item— (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T
LAC		214-218, 68, The American Society of Human Genetics.	1
4e		ZHOU, Guo-Hua, et al., Quantitative detection of single nucleotide polymorphisms for a pooled sample by a bioluminometric assay coupled with modified primer extension reactions (BAMPER), Nucleic Acids Research, (2001) 1-11, 29(19 e93), Oxford University Press.	

EXAMINER Suid Clar DATE CONSIDERED 3/16/05